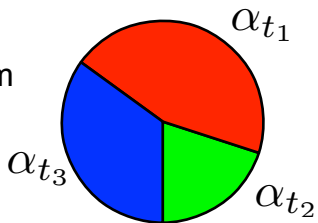
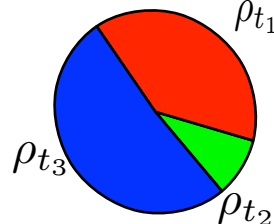


probabilities
of selecting
fragments from
each of the
transcripts



$$\rho_t = \frac{\frac{\alpha_t}{l_t}}{\sum_r \frac{\alpha_r}{l_r}}$$

$$\alpha_t = \frac{\rho_t l_t}{\sum_r \rho_r l_r}$$



relative
transcript
abundances

two alignments of a fragment f

